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Pastorek, Jaromir

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gaggatctac ctggagagga ggatctacct gaagttaagc ctaaatcaga 350

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<220>
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tggagagaaa ataaaaaggg tgcaaaagga gagaggtgag ctggatgaga tgggagagaa 240

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<212> DNA
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<222> (1)..(131)
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<400> 41
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cacgttggga ggctgaggtg ggagaatggt ttgagcccag gagttcaaga caaggcgggg 180

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<210> 46
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<220>
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<222> (1)..(114)
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<400> 46
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gtgtctgtca ttggtggtca cagcccgcct ctcacatctc ctttttctct ccag 114

<210> 47
<211> 617
<212> DNA
<213> HUMAN

<220>
<221> intron
<222> (1)..(617)
<223> 9th MN intron

<400> 47
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agggctgctc aggaccgcct ctgctccctc tccttttctg cagaacagac cccaacccca 120

atattagaga ggcagatcat ggtggggatt cccccattgt ccccagaggc taattgatta 180

gaatgaagct tgagaaatct cccagcatcc ctctcgcaaa agaatccccc ccccttttt 240

taaagatagg gtctcactct gtttgcccca ggctggggtg ttgtggcacg atcatagctc 300

actgcagcct cgaactccta ggctcaggca atcctttcac cttagcttct caaagcactg 360

ggactgtagg catgagccac tgtgcctggc cccaaacggc ccttttactt ggcttttagg 420

aagcaaaaac ggtgcttatc ttacccccttc tcgtgtatcc accctcatcc cttggctggc 480

ctcttctgga gactgaggca ctatggggct gcctgagaac tcggggcagg ggtggtggag 540

tgcactgagg caggtgttga ggaactctgc agacccctct tccttcccaa agcagccctc 600

tctgctctcc atcgcag 617

<210> 48
<211> 130
<212> DNA
<213> HUMAN

<220>
<221> intron
<222> (1)..(130)
<223> 10th MN intron

<400> 48
gtattacact gacccttttct tcaggcacaa gcttccccca cccttgtgga gtcacttcat 60

gcaaagcgca tgcaaatgag ctgctcctgg gccagttttc tgattagcct ttcctgttgt 120

gtacacacag 130

<210> 49
<211> 1401
<212> DNA
<213> HUMAN

<400> 49
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cctccacctc ccgggttcaa gtgattctcc tgcctcagcc tctagccaag tagctgcgat 120

tacaggcatg cgccaccacg cccggctaat ttttgtattt ttagtagaga cggggtttcg 180

ccatgttggt caggctggtc tcgaactcct gatctcaggt gatccaacca ccctggcctc 240

ccaaagtgct gggattatag gcgtgagcca cagcgcctgg cctgaagcag ccactcactt 300

ttacagaccc taagacaatg attgcaagct ggtaggattg ctgtttggcc cacccagctg 360

cggtgttgag tttgggtgcg gtctcctgtg ctttgcacct ggcccgctta aggcatttgt 420

tacccgtaat gctcctgtaa ggcatctgcg tttgtgacat cgttttggtc gccaggaagg 480

gattggggct ctaagcttga gcggttcatc cttttcattt atacagggga tgaccagagt 540

cattggcgct atggaggtga gacacccacc cgctgcacag acccaatctg ggaacccagc 600

tctgtggatc tcccctacag ccgtccctga acactggtcc cgggcgtccc acccgccgcc 660

caccgtccca ccccctcacc ttttctaccc gggttcccta agttcctgac ctaggcgtca 720

gacttcctca ctatactctc ccaccccagg cgacccgccc tggccccggg tgtccccagc 780

ctgcgcgggc cgcttccagt ccccggtgga tatccgcccc cagctcgccg ccttctgccc 840

ggccctgcgc cccctggaac tcctgggctt ccagctcccg ccgctcccag aactgcgcct 900

gcgcaacaat ggccacagtg gtgagggggt ctccccgccg agacttgggg atggggcggg 960

gcgcagggaa gggaaccgtc gcgcagtgcc tgcccggggg ttgggctggc cctaccgggc 1020

ggggccggct cacttgcctc tccctacgca gtgcaactga ccctgcctcc tgggctagag 1080

atggctctgg gtcccgggcg ggagtaccgg gctctgcagc tgcatctgca ctgggggggct 1140

gcaggtcgtc cgggctcgga gcacactgtg gaaggccacc gtttccctgc cgaggtgagc 1200

gcggactggc cgagaagggg caaaggagcg gggcggacgg gggccagaga cgtggccctc 1260

tcctaccctc gtgtcctttt cagatccacg tggttcacct cagcaccgcc tttgccagag 1320

ttgacgaggc cttggggcgc ccgggaggcc tggccgtgtt ggccgccttt ctggaggtac 1380

cagatcctgg acacccccta c 1401

<210> 50
<211> 59
<212> PRT
<213> HUMAN

<400> 50
Ser Ser Gly Glu Asp Asp Pro Leu Gly Glu Glu Asp Leu Pro Ser Glu
1 5 10 15

Glu Asp Ser Pro Arg Glu Glu Asp Pro Pro Gly Glu Glu Asp Leu Pro
20 25 30

Gly Glu Glu Asp Leu Pro Gly Glu Glu Asp Leu Pro Glu Val Lys Pro
35 40 45

Lys Ser Glu Glu Glu Gly Ser Leu Lys Leu Glu
50 55

<210> 51
<211> 257
<212> PRT
<213> HUMAN

<400> 51

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Gly Asp Asp Gln Ser His Trp Arg Tyr Gly Gly Asp Pro Pro Trp Pro
 1               5                  10                  15

Arg Val Ser Pro Ala Cys Ala Gly Arg Phe Gln Ser Pro Val Asp Ile
            20                  25                  30

Arg Pro Gln Leu Ala Ala Phe Cys Pro Ala Leu Arg Pro Leu Glu Leu
        35                  40                  45

Leu Gly Phe Gln Leu Pro Pro Leu Pro Glu Leu Arg Leu Arg Asn Asn
    50                  55                  60

Gly His Ser Val Gln Leu Thr Leu Pro Pro Gly Leu Glu Met Ala Leu
65                  70                  75                  80

Gly Pro Gly Arg Glu Tyr Arg Ala Leu Gln Leu His Leu His Trp Gly
                85                  90                  95

Ala Ala Gly Arg Pro Gly Ser Glu His Thr Val Glu Gly His Arg Phe
            100                 105                 110

Pro Ala Glu Ile His Val Val His Leu Ser Thr Ala Phe Ala Arg Val
        115                 120                 125

Asp Glu Ala Leu Gly Arg Pro Gly Gly Leu Ala Val Leu Ala Ala Phe
    130                 135                 140

Leu Glu Glu Gly Pro Glu Glu Asn Ser Ala Tyr Glu Gln Leu Leu Ser
145                 150                 155                 160

Arg Leu Glu Glu Ile Ala Glu Glu Gly Ser Glu Thr Gln Val Pro Gly
                165                 170                 175

Leu Asp Ile Ser Ala Leu Leu Pro Ser Asp Phe Ser Arg Tyr Phe Gln
            180                 185                 190

Tyr Glu Gly Ser Leu Thr Thr Pro Pro Cys Ala Gln Gly Val Ile Trp
        195                 200                 205

Thr Val Phe Asn Gln Thr Val Met Leu Ser Ala Lys Gln Leu His Thr
    210                 215                 220

Leu Ser Asp Thr Leu Trp Gly Pro Gly Asp Ser Arg Leu Gln Leu Asn
225                 230                 235                 240

Phe Arg Ala Thr Gln Pro Leu Asn Gly Arg Val Ile Glu Ala Ser Phe
                245                 250                 255

Pro
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<210> 52
<211> 20
<212> PRT
<213> HUMAN

<400> 52

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Ile Leu Ala Leu Val Phe Gly Leu Leu Phe Ala Val Thr Ser Val Ala
  1               5                  10                  15

Phe Leu Val Gln
             20
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<210> 53
<211> 25
<212> PRT
<213> HUMAN

<400> 53

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Met Arg Arg Gln His Arg Arg Gly Thr Lys Gly Gly Val Ser Tyr Arg
  1               5                  10                  15

Pro Ala Glu Val Ala Glu Thr Gly Ala
             20                  25
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<210> 54
<211> 59
<212> PRT
<213> HUMAN

<400> 54

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Ser Ala Ser Glu Glu Pro Ser Pro Ser Glu Val Pro Phe Pro Ser Glu
  1               5                  10                  15

Glu Pro Ser Pro Ser Glu Glu Pro Phe Pro Ser Val Arg Pro Phe Pro
             20                  25                  30

Ser Val Val Leu Phe Pro Ser Glu Glu Pro Phe Pro Ser Lys Glu Pro
         35                  40                  45

Ser Pro Ser Glu Glu Pro Ser Ala Ser Glu Glu
     50                  55
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<210> 55
<211> 470
<212> RNA
<213> HUMAN

<400> 55

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cauggccccg auaaccuucu gccugugcac acaccugccc cucacuccac ccccauccua 60

gcuuugguau gggggagagg gcacagggcc agacaaaccu gugagacuuu ggcuccaucu 120

cugcaaaagg gcgcucugug agucagccug cuccccucca ggcuugcucc ucccccaccc 180

agcucucguu uccaaugcac guacagcccg uacacaccgu gugcugggac accccacagu 240

cagccgcaug gcuccccugu gccccagccc cuggcucccu cuguugaucc cggccccugc 300

uccaggccuc acugugcaac ugcugcuguc acugcugcuu cuggugccug uccaucccca 360

gagguugccc cggaugcagg aggauucccc cuugggagga ggcucuucug gggaagauga 420
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cccacugggc gaggaggauc ugcccaguga agaggauuca cccagagagg 470

<210> 56
<211> 292
<212> DNA
<213> HUMAN

<400> 56
gttttttga gacggagtct tgcatctgtc atgcccaggc tggagtagca gtggtgccat 60

ctcggctcac tgcaagctcc acctcccgag ttcacgccat tttcctgcct cagcctcccg 120

agtagctggg actacaggcg cccgccacca tgcccggcta atttttttgta tttttggtag 180

agacggggtt tcaccgtgtt agccagaatg gtctcgatct cctgacttcg tgatccaccc 240

gcctcggcct cccaaagttc tgggattaca ggtgtgagcc accgcacctg gc 292

<210> 57
<211> 262
<212> DNA
<213> HUMAN

<400> 57
tttctttttt gagacagggt cttgctctgt cacccaggcc agagtgcaat ggtacagtct 60

cagctcactg cagcctcaac cgcctcggct caaaccatca tcccatttca gcctcctgag 120

tagctgggac tacaggcaca tgccattaca cctggctaat ttttttgtat ttctagtaga 180

gacagggttt ggccatgttg cccgggctgg tctcgaactc ctggactcaa gcaatccacc 240

cacctcagcc tcccaaaatg ag 262

<210> 58
<211> 2501
<212> DNA
<213> HUMAN

<220>
<221> misc_feature
<222> (1)..(2501)

<400> 58
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agggttaaat ggattaaggg cggtgcaaga tgtgctttgt taaacagatg cttgaaggca 120

gcatgctcgt taagagtcat caccaatccc taatctcaag taatcaggga cacaaacact 180

gcggaaggcc gcagggtcct ctgcctagga aaaccagaga cctttgttca cttgtttatc 240

tgaccttccc tccactattg tccatgaccc tgccaaatcc ccctctgtga gaaacaccca 300

agaattatca ataaaaaaat aaatttaaaa aaaaaataca aaaaaaaaaa aaaaaaaaaa 360

aaaagactta cgaatagtta ttgataaatg aatagctatt ggtaaagcca agtaaatgat 420

catattcaaa accagacggc catcatcaca gctcaagtct acctgatttg atctctttat 480

cattgtcatt ctttggattc actagattag tcatcatcct caaaattctc ccccaagttc 540

taattacgtt ccaaacattt aggggttaca tgaagcttga acctactacc ttctttgctt 600

ttgagccatg agttgtagga atgatgagtt tacaccttac atgctgggga ttaatttaaa 660

ctttacctct aagtcagttg ggtagccttt ggcttatttt tgtagctaat tttgtagtta 720

atggatgcac tgtgaatctt gctatgatag ttttcctcca cactttgcca ctaggggtag 780

gtaggtactc agttttcagt aattgcttac ctaagaccct aagccctatt tctcttgtac 840

tggcctttat ctgtaatatg ggcatattta atacaatata atttttggag ttttttttgtt 900

tgtttgtttg tttgtttttt tgagacggag tcttgcatct gtcatgccca ggctggagta 960

gcagtggtgc catctcggct cactgcaagc tccacctccc gagttcacgc cattttcctg 1020

cctcagcctc ccgagtagct gggactacag gcgcccgcca ccatgcccgg ctaatttttt 1080

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ctggccaatt ttttgagtct tttaaagtaa aaatatgtct tgtaagctgg taactatggt 1260

acatttcctt ttattaatgt ggtgctgacg gtcatatagg ttcttttgag tttggcatgc 1320

atatgctact ttttgcagtc ctttcattac atttttctct cttcatttga agagcatgtt 1380

atatctttta gcttcacttg gcttaaaagg ttctctcatt agcctaacac agtgtcattg 1440

ttggtaccac ttggatcata agtggaaaaa cagtcaagaa attgcacagt aatacttgtt 1500

tgtaagaggg atgattcagg tgaatctgac actaagaaac tcccctacct gaggtctgag 1560

attcctctga cattgctgta tataggcttt tcctttgaca gcctgtgact gcggactatt 1620

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tgcatcaagt gagaacatat aatgtctgca tgtttccata tttcaggaat gtttgcttgt 1740

gttttatgct tttatataga cagggaaact tgttcctcag tgacccaaaa gaggtgggaa 1800

ttgttattgg atatcatcat tggcccacgc tttctgacct tggaaacaat taagggttca 1860

taatctcaat tctgtcagaa ttggtacaag aaatagctgc tatgtttctt gacattccac 1920

ttggtaggaa ataagaatgt gaaactcttc agttggtgtg tgtccctngt ttttttgcaa 1980

tttccttctt actgtgttaa aaaaaagtat gatcttgctc tgagaggtga ggcattctta 2040

atcatgatct ttaaagatca ataatataat cctttcaagg attatgtctt tattataata 2100

aagataattt gtctttaaca gaatcaataa tataatccct taaaggatta tatctttgct 2160

gggcgcagtg gctcacacct gtaatcccag cactttgggt ggccaaggtg gaaggatcaa 2220

atttgcctac ttctatatta tcttctaaag cagaattcat ctctcttccc tcaatatgat 2280

gatattgaca gggtttgccc tcactcacta gattgtgagc tcctgctcag ggcaggtagc 2340

gtttttttgtt tttgtttttg ttttttctttt ttgagacagg gtcttgctct gtcacccagg 2400

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catcccattt cagcctcctg agtagctggg actacaggca c 2501

<210> 59
<211> 292
<212> DNA
<213> HUMAN

<220>
<221> misc_feature
<222> (1)

<400> 59
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tcggctcact gcaagctcca cctcccgagt tcacgccatt ttcctgcctc agcctcccga 120

gtagctggga ctacaggcgc ccgccaccat gcccggctaa ttttttgtat ttttggtaga 180

gacggggttt caccgtgtta gccagaatgg tctcgatctc ctgacttcgt gatccacccg 240

cctcggcctc ccaaagttct gggattacag gtgtgagcca ccgcacctgg cc 292

<210> 60
<211> 262
<212> DNA
<213> HUMAN

<400> 60
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agctcactgc agcctcaacc gcctcggctc aaaccatcat cccatttcag cctcctgagt 120

agctgggact acaggcacat gccattacac ctggctaatt tttttgtatt tctagtagag 180

acagggtttg gccatgttgc ccgggctggt ctcgaactcc tggactcaag caatccaccc 240

acctcagcct cccaaaatga gg 262

<210> 61
<211> 294
<212> DNA
<213> HUMAN

<400> 61
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cggctcactg caacctccac ctcccgggtt caagtgattc tcctgcctca gcctctagcc 120

aagtagctgc gattacaggc atgcgccacc acgcccggct aatttttgta tttttagtag 180

agacggggtt tcgccatgtt ggtcaggctg gtctcgaact cctgatctca ggtgatccaa 240

ccaccctggc ctcccaaagt gctgggatta taggcgtgag ccacagcgcc tggc 294

<210> 62
<211> 276
<212> DNA
<213> HUMAN

<400> 62
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tccgcctccc gggttcaagg gattctcctg cctcagcttc ctgagtagct ggggttacag 120

gtgtgtgcca ccatgcccag ctaatttttt tttgtatttt tagtagacag ggtttcacca 180

tgttggtcag gctggtctca aactcctggc ctcaagtgat ccgcctgact cagcctacca 240

aagtgctgat tacaagtgtg agccaccgtg cccagc 276

<210> 63
<211> 289
<212> DNA
<213> HUMAN

<400> 63
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tcacgaggtc aagagatcaa gaccatcctg gccaacatgg tgaaacccca tctctactaa 120

aaatacgaaa aaatagccag gcgtggtggc gggtgcctgt aatcccagct actcgggagg 180

ctgaggcagg agaatggcat gaacccggga ggcagaagtt gcagtgagcc gagatcgtgc 240

cactgcactc cagcctgggc aacagagcga gactcttgtc tcaaaaaaa 289

<210> 64
<211> 298
<212> DNA
<213> HUMAN

<400> 64
aggctgggct ctgtggctta cgcctataat cccaccacgt tgggaggctg aggtgggaga 60

atggtttgag cccaggagtt caagacaagg cggggcaaca tagtgtgacc ccatctctac 120

caaaaaaacc ccaacaaaac caaaaatagc cgggcatggt ggtatgcggc ctagtcccag 180

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gctatgatcc caccactgcc taccatcttt aggatacatt tatttattta taaaagaa 298

<210> 65
<211> 105
<212> DNA
<213> HUMAN

<400> 65
ttttttacat ctttagtaga gacagggttt caccatattg gccaggctgc tctcaaactc 60

ctgaccttgt gatccaccag cctcggcctc ccaaagtgct gggat 105

<210> 66
<211> 83
<212> DNA
<213> HUMAN

<400> 66
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aggcatgagc cactgtgcct ggc 83

<210> 67
<211> 11
<212> DNA
<213> HUMAN

<400> 67
agaaggtaag t 11

<210> 68
<211> 11
<212> DNA
<213> HUMAN

<400> 68
tggaggtgag a 11

<210> 69
<211> 11
<212> DNA
<213> HUMAN

<400> 69
cagtcgtgag g 11

<210> 70
<211> 11
<212> DNA
<213> HUMAN

<400> 70
ccgaggtgag c 11

<210> 71
<211> 11
<212> DNA
<213> HUMAN

<400> 71
tggaggtacc a 11

<210> 72
<211> 11
<212> DNA
<213> HUMAN

<400> 72
ggaaggtcag t 11

<210> 73
<211> 11
<212> DNA
<213> HUMAN

<400> 73
agcaggtggg c 11

<210> 74
<211> 11
<212> DNA
<213> HUMAN

<400> 74
gccaggtaca g 11

<210> 75
<211> 11
<212> DNA
<213> HUMAN

<400> 75
tgctggtgag t 11

<210> 76
<211> 11
<212> DNA
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<400> 76
cacaggtatt a 11

<210> 77
<211> 11
<212> DNA
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<400> 77
atacagggga t 11

<210> 78
<211> 11
<212> DNA
<213> HUMAN

<400> 78
ccccaggcga c 11

<210> 79
<211> 11
<212> DNA
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<400> 79
acgcagtgca a 11

<210> 80
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<400> 80
tttcagatcc a 11

<210> 81
<211> 11
<212> DNA
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<400> 81
ccccaggagg g 11

<210> 82
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<400> 82
tcacaggctc a 11

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ccctagctcc a 11

<210> 84
<211> 11
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<213> HUMAN

<400> 84
ctccagtcca g 11

<210> 85
<211> 12
<212> DNA
<213> HUMAN

<400> 85
tcgcaggtga ca 12

<210> 86
<211> 11
<212> DNA
<213> HUMAN

<400> 86
acacagaagg g 11

<210> 87
<211> 377
<212> PRT
<213> HUMAN

<400> 87

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Gln Arg Leu Pro Arg Met Gln Glu Asp Ser Pro Leu Gly Gly Gly Ser
  1               5                  10                  15

Ser Gly Glu Asp Asp Pro Leu Gly Glu Glu Asp Leu Pro Ser Glu Glu
             20                  25                  30

Asp Ser Pro Arg Glu Glu Asp Pro Pro Gly Glu Glu Asp Leu Pro Gly
         35                  40                  45

Glu Glu Asp Leu Pro Gly Glu Glu Asp Leu Pro Glu Val Lys Pro Lys
     50                  55                  60

Ser Glu Glu Glu Gly Ser Leu Lys Leu Glu Asp Leu Pro Thr Val Glu
 65                  70                  75                  80

Ala Pro Gly Asp Pro Gln Glu Pro Gln Asn Asn Ala His Arg Asp Lys
                 85                  90                  95

Glu Gly Asp Asp Gln Ser His Trp Arg Tyr Gly Gly Asp Pro Pro Trp
            100                 105                 110
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Pro Arg Val Ser Pro Ala Cys Ala Gly Arg Phe Gln Ser Pro Val Asp
115 120 125

Ile Arg Pro Gln Leu Ala Ala Phe Cys Pro Ala Leu Arg Pro Leu Glu
130 135 140

Leu Leu Gly Phe Gln Leu Pro Pro Leu Pro Glu Leu Arg Leu Arg Asn
145 150 155 160

Asn Gly His Ser Val Gln Leu Thr Leu Pro Pro Gly Leu Glu Met Ala
165 170 175

Leu Gly Pro Gly Arg Glu Tyr Arg Ala Leu Gln Leu His Leu His Trp
180 185 190

Gly Ala Ala Gly Arg Pro Gly Ser Glu His Thr Val Glu Gly His Arg
195 200 205

Phe Pro Ala Glu Ile His Val Val His Leu Ser Thr Ala Phe Ala Arg
210 215 220

Val Asp Glu Ala Leu Gly Arg Pro Gly Gly Leu Ala Val Leu Ala Ala
225 230 235 240

Phe Leu Glu Glu Gly Pro Glu Glu Asn Ser Ala Tyr Glu Gln Leu Leu
245 250 255

Ser Arg Leu Glu Glu Ile Ala Glu Glu Gly Ser Glu Thr Gln Val Pro
260 265 270

Gly Leu Asp Ile Ser Ala Leu Leu Pro Ser Asp Phe Ser Arg Tyr Phe
275 280 285

Gln Tyr Glu Gly Ser Leu Thr Thr Pro Pro Cys Ala Gln Gly Val Ile
290 295 300

Trp Thr Val Phe Asn Gln Thr Val Met Leu Ser Ala Lys Gln Leu His
305 310 315 320

Thr Leu Ser Asp Thr Leu Trp Gly Pro Gly Asp Ser Arg Leu Gln Leu
325 330 335

Asn Phe Arg Ala Thr Gln Pro Leu Asn Gly Arg Val Ile Glu Ala Ser
340 345 350

Phe Pro Ala Gly Val Asp Ser Ser Pro Arg Ala Ala Glu Pro Val Gln
355 360 365

Leu Asn Ser Cys Leu Ala Ala Gly Asp
370 375

<210> 88
<211> 34
<212> DNA
<213> HUMAN

<400> 88

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<210> 89
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<212> DNA
<213> HUMAN

<400> 89
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<210> 90
<211> 3532
<212> DNA
<213> HUMAN

<220>
<221> misc_feature
<222> (1)..(3532)

<400> 90
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agggttaaat ggattaaggg cggtgcaaga tgtgctttgt taaacagatg cttgaaggca 120

gcatgctcgt taagagtcat caccaatccc taatctcaag taatcaggga cacaaacact 180

gcggaaggcc gcagggtcct ctgcctagga aaaccagaga cctttgttca cttgtttatc 240

tgaccttccc tccactattg tccatgaccc tgccaaatcc ccctctgtga gaaacaccca 300

agaattatca ataaaaaaat aaatttaaaa aaaaaataca aaaaaaaaaa aaaaaaaaaa 360

aaaagactta cgaatagtta ttgataaatg aatagctatt ggtaaagcca agtaaatgat 420

catattcaaa accagacggc catcatcaca gctcaagtct acctgatttg atctctttat 480

cattgtcatt ctttggattc actagattag tcatcatcct caaaattctc ccccaagttc 540

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